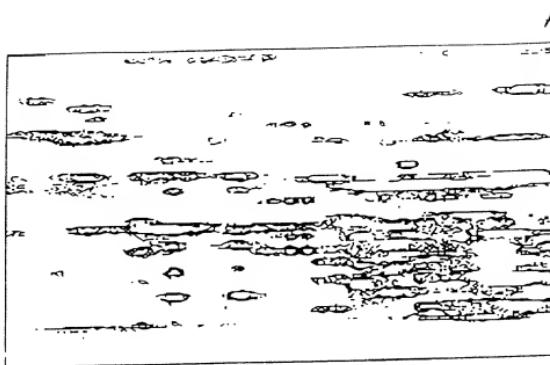


FIGURE 1A

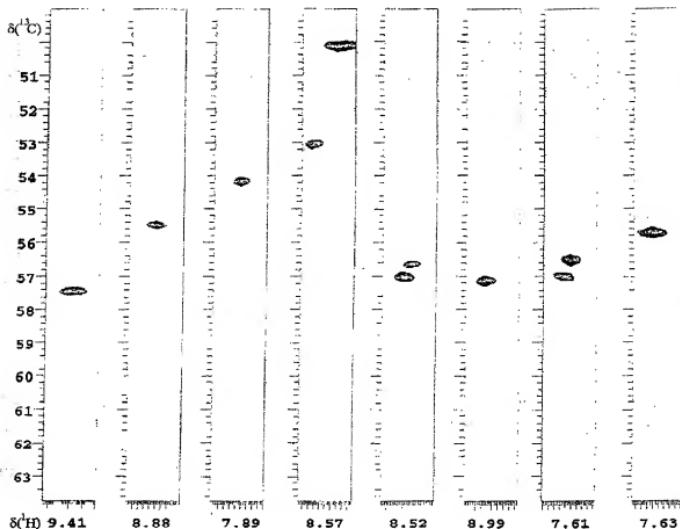


09983820.102204

TOEOT: 02032565

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Figure 1B of 13

FIGURE 1B



Dipolar Couplings That Depend Only on $\phi(i)$ and $\psi(i)$



Search ϕ and ψ Until Measured Couplings = Theoretical Couplings

Packing Secondary Structural Elements

FIGURE 3

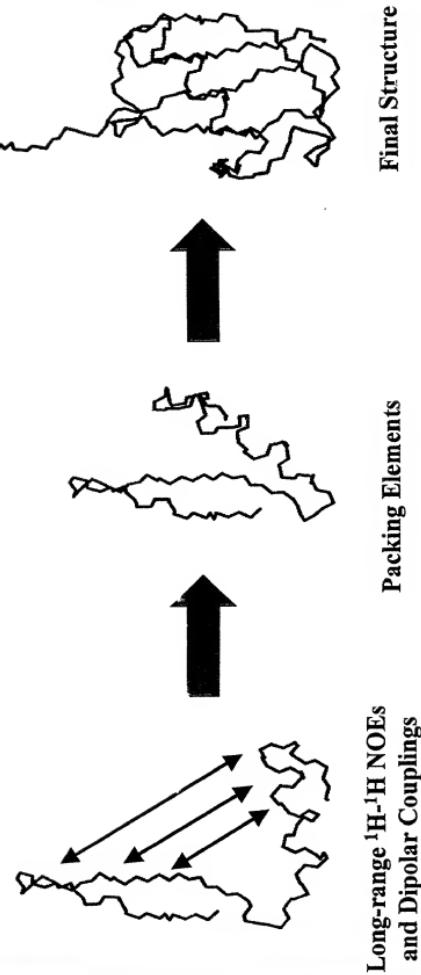
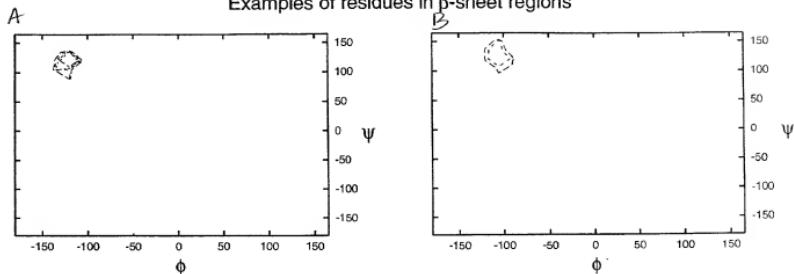


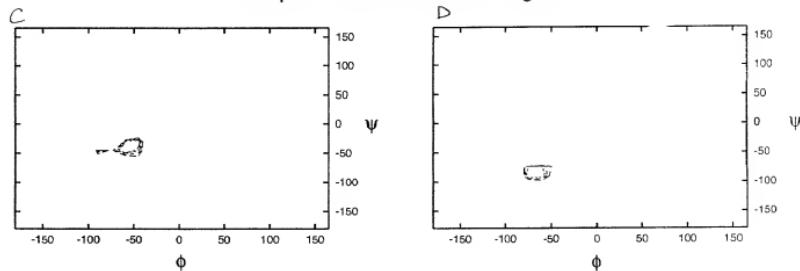
FIGURE 4

Examples of residues in β -sheet regions

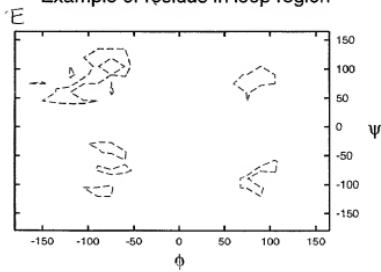


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Examples of residues in α -helix regions



Example of residue in loop region



0208604, 0208604

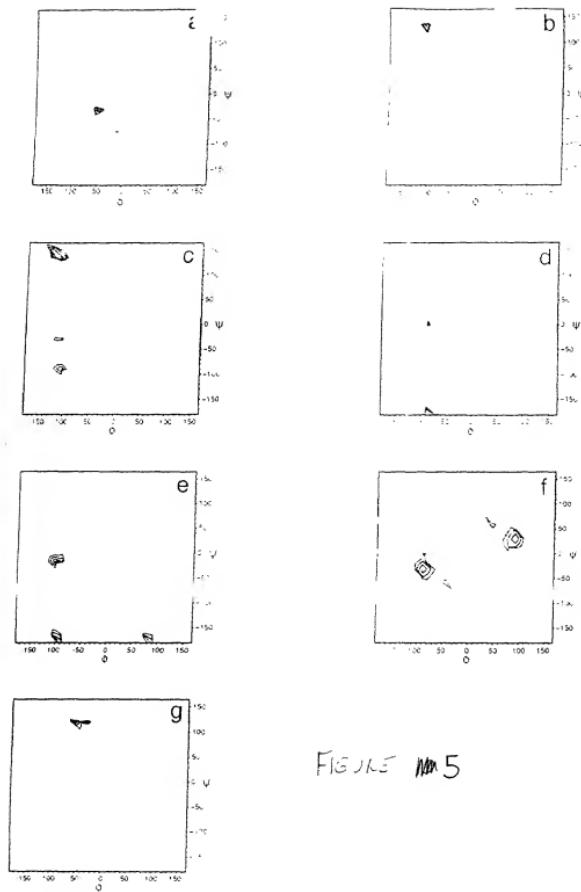


FIGURE NO 5

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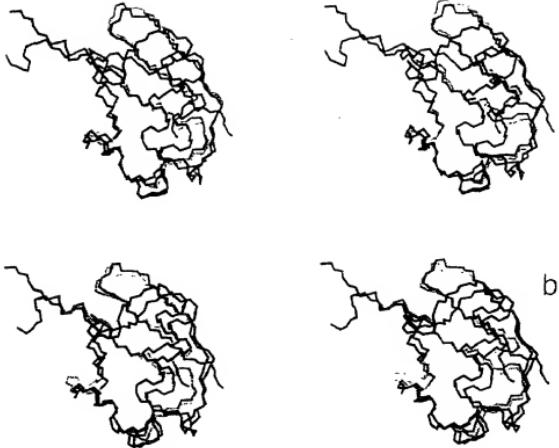


FIGURE 6

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FIGURE 7

Generate Linear
amino-acid chain

Calculate ϕ, ψ angles
for each peptide pair
using experimental
residual dipolar couplings

Fold Linear sequence
with dihedral angle
and backbone NOE
restraints

Refine structure
using NOE and
dipolar coupling
restraints

102201-0003333301

Dipolar Couplings - Powerful Structural Constraints

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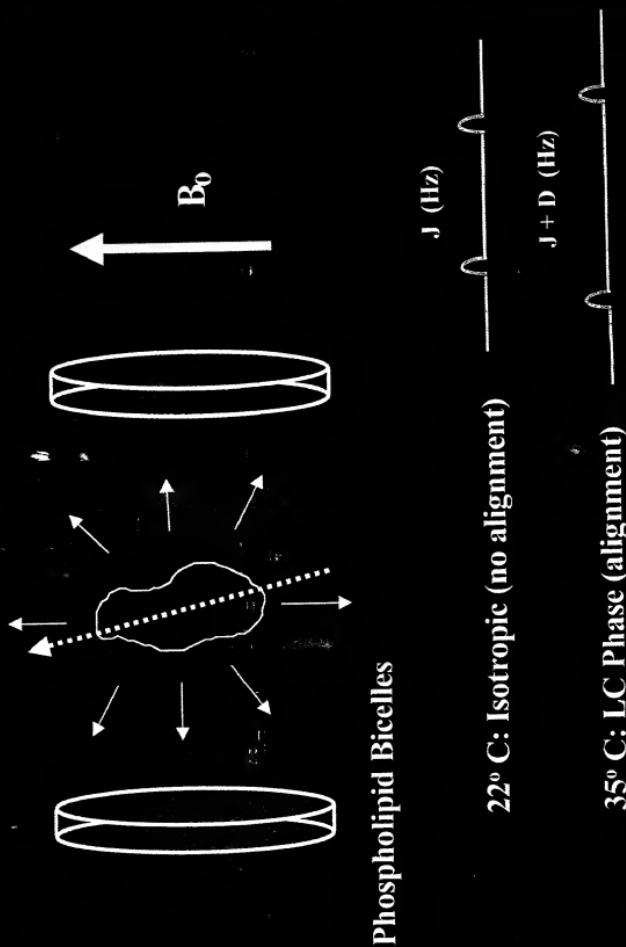


$$D \propto (3 \cos^2 \theta - 1) / r^3$$



FIGURE 9

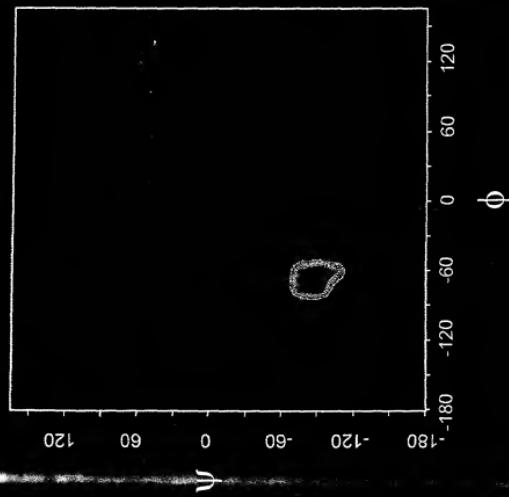
Measurement of Dipolar Couplings Requires a Weakly Aligned Molecule



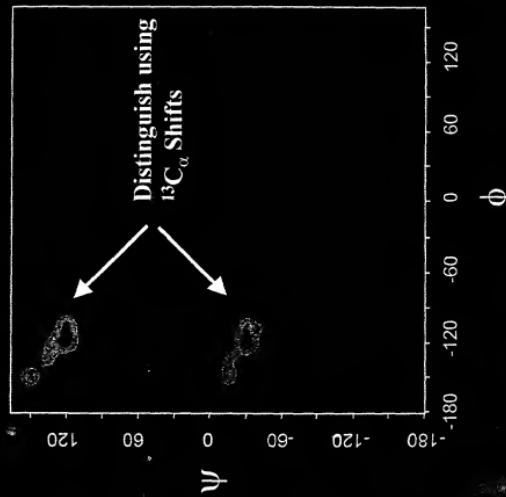
ϕ, ψ Mapping Using Residual Dipolar Couplings

FIGURE 10

Val 26



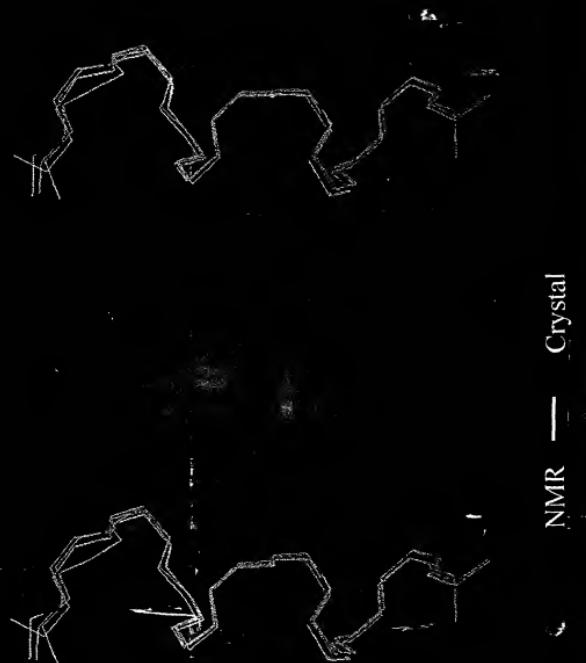
Ile 33



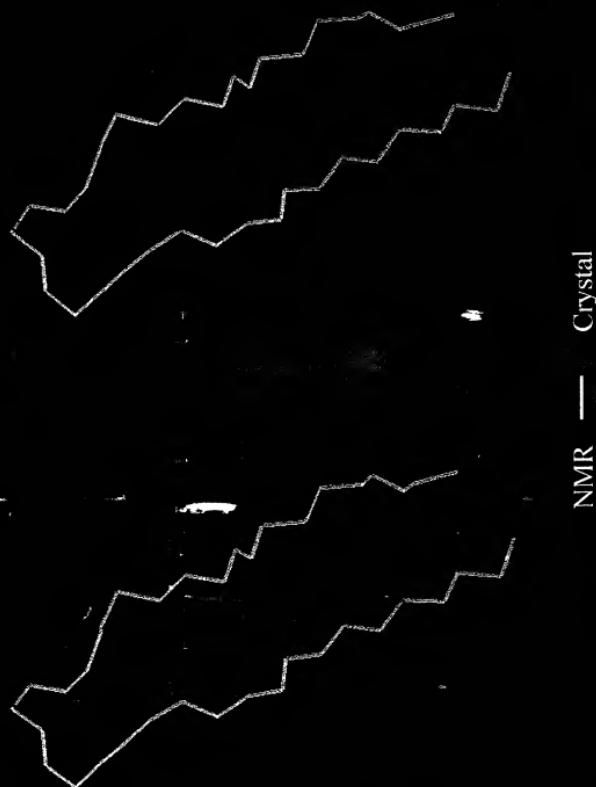
Distinguish using
 $^{13}\text{C}_\alpha$ Shifts

16
FIGURE 11. α-helices of Ubiquitin.

NMR vs. Crystal Structure of α -helix (24-34) Ubiquitin



13
FIGURE 12
NMR vs. Crystal Structure of β -sheet (3-15) Ubiquitin



A 4 0 *

FIGURE 12

Crystal Structure vs. NMR Global Fold - Ubiquitin

